

# PCR protocol of XynB, ArfB and ManA1

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In previous step, we have successfully isolated pure culture of ***Clostridium stecorarium*** and cultivated it on solid medium in our lab. Then colony PCR was conducted to amplify the genes of interest from *C. stecorarium*.

The genes of interest include  **$\alpha$ -L-arabinofuranosidase gene (ArfB)**, **endo-1,4-beta-xylanase gene (XynB)** and **mannan endo-1,4-beta-mannosidase gene(ManA1)**, three candidates for biobleaching.

In the following PCR reactions, the genome from ***E.coli* (DH5 $\alpha$ )** other than *C. stecorarium* was used as template in the negative control groups.

We picked single colony from the solid plate and diluted it in **10  $\mu$ l LB liquid medium**, then used **1 $\mu$ l** as template for colony PCR.

The recipe of the reaction mixture is shown as below:

Template	1 $\mu$ l
Forward primer	1 $\mu$ l
Reverse primer	1 $\mu$ l
Taq master mix (Tiangen)	12.5 $\mu$ l
H2O	4.5 $\mu$ l
Total	20 $\mu$ l

According to the results of gradient PCR (data not shown), the proximal annealing temperatures for the three genes were separately determined: **53.7°C for XynB** , **51°C for ArfB**, and **53.1°C for ManA1**.

The PCR program for gene amplification is carried out as below:

Temperature(°C)			Time(s)
XynB	ArfB	ManA1	
94	94	94	30
53.7	51	51.1	30
72	72	72	120
72	72	72	600

35 repeats

(Before PCR, the bacteria were heated at 98°C for 20 minutes in order to be lysed fully. )

Then, the PCR products (the result is shown as below) was detected by **1% agrose** gel electrophoresis . The PCR products were consistent with the expected size according to the whole genome sequencing for ***Clostridium stercorarium* DSM8532 (Poehlein et al, 2013)**, which the fragments of **XynB, ArfB and ManA1** were respectively **1164bp, 1473bp, and 1530bp**.

